

71914.ST25.txt SEQUENCE LISTING

COUNCIL OF SCENTIFIC AND INDUSTRIAL RESEARCH <110>

<120> STABLE GENE VARIANTS OF LIPASES

<130> 71914

<140> us 10/768,951

2004-01-29 <141>

22 <160>

<170> PatentIn version 3.3

<210>

181 <211>

<212> PRT

Bacillus subtilis <213>

<220>

AMINO ACIDS

<221> <222> (1)..(181)

<223> enzyme sequence

<400>

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 5 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 55 60

Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly
100 105 110

Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser

Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Asn Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175

Gly Gln Asn Thr Asn 180

<210> 2 181 <211>

<212> PRT

Bacillus subtilis <213>

<220>

Amino acid (1)..(181) <221> <222>

Protein sequence <223>

<400> 2

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105

Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser

Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 135 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu Page 2

150

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175

Gly Gln Asn Thr Asn 180

<210>

<211> 181 <212> PRT

Bacillus subtilis <213>

<220>

<221>

<222>

Amino acid (1)..(181) Protein sequence

<400> 3

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Val Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105

Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser 115 120 125

Ile Tyr Ser Ser Asp Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu
130 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly
165 170 175

Gly Gln Asn Thr Asn

<210>

181 <211>

<212> **PRT** Bacillus subtilis

<220>

<221>

Amino acid (1)..(181) <222>

Protein Sequence <223>

<400>

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 5 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu
50 60

Asp Glu Thr Gly Thr Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105 110

Lys Ala Pro Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser

Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 135 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly Page 4

Gly Gln Asn Thr Asn 180

<210> 5 <211> 181

<211> 101 <212> PRT

<213> Bacillus subtilis

<220>

<221> Amino acid <222> (1)..(181)

<223> Protein Sequence

<400> 5

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105 110

Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser 115 120 125

Ile Tyr Ser Ser Asp Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 135 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175

Gly Gln Asn Thr Asn 180

<210> 6

<211> 181 <212> PRT

Bacillus subtilis

<220>

<221>

<222>

Amino acid (1)..(181) Protein sequence <223>

<400>

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 5 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 . 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105 110

Lys Ala Pro Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser 115 120 125

Ile Tyr Ser Ser Asp Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 135 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly
165 170 175

Gly Gln Asn Thr Asn

```
7
181
<210>
<211>
<212>
        PRT
        Bacillus subtilis
<220>
<221>
        Amino acid
<222>
        (1)..(181)
<223>
        Protein sequence
<400>
Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala
1 5 10 15
Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp
20 25 30
Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45
Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60
Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80
Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys
85 90 95
Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105 110
Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser 115 120 125 .
Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu
130 135 140
Asp Gly Ala Ser Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 160
Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly
165 170 175
Gly Gln Asn Thr Asn
180
```

Page 7

```
<210>
        181
<211>
<212>
        PRT
        Bacillus subtilis
<220>
<221>
<222>
        Amino acid (1)..(181)
        Protein sequence
<400>
Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala
1 10 15
Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp
20 25 30
Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45
Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60
Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80
Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys
85 90 95
Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105 110
Lys Ala Pro Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser
115 120 125
Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu
130 135 140
Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu
145 150 155 160
Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175
Gly Gln Asn Thr Asn
```

<210> 9

180

<211> 181

<212> PRT

Bacillus subtilis

<220>

Amino acid (1)..(181) <221>

<222>

<223> Protein Sequence

<400>

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Ala Lys Lys Ala Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105

Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser

Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly
165 170 175

Gly Gln Asn Thr Asn 180

<210> 10

<211> 181 **PRT**

<213> Bacillus subtilis

<220> .

<221> <222> Amino acid (1)..(181)

<223> Protein Sequence

<400> 10

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 5 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105 110

Lys Ala Leu Pro Gly Thr Asp Pro Asn Gln Lys Ile Leu Tyr Thr Ser 115 120 125

Ile Tyr Ser Ser Ala Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 135 140

Val Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175

Gly Gln Asn Thr Asn 180

<210> 11

<211> 181 PRT

<212> Bacillus subtilis

```
<220>
        Amino acid (1)..(181)
<221>
<222>
        Protein sequence
<400>
        11
Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala
1 10 15
Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30
Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45
Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60
Asp Glu Thr Gly Val Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80
Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys
85 90 95
Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly
100 105 110
Lys Ala Leu Pro Gly Thr Asp Pro Asp Gln Lys Ile Leu Tyr Thr Ser
115 120 125
Ile Tyr Ser Ser Asp Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu
130 135 140
Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu
145 150 155 160
Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175
Gly Gln Asn Thr Asn
              180
```

<220>

<210> 12

<211> 181

<212> PRT

<213> Bacillus subtilis

<221>

<222>

Amino acid (1)..(181) Protein sequence

<400> 12

Met Ala Glu His Asn Pro Val Val Met Val His Gly Ile Gly Gly Ala 1 10 15

Ser Phe Asn Phe Ala Gly Ile Lys Ser Tyr Leu Val Ser Gln Gly Trp 20 25 30

Ser Arg Asp Lys Leu Tyr Ala Val Asp Phe Trp Asp Lys Thr Gly Thr 35 40 45

Asn Tyr Asn Asn Gly Pro Val Leu Ser Arg Phe Val Gln Lys Val Leu 50 60

Asp Glu Thr Gly Ala Lys Lys Val Asp Ile Val Ala His Ser Met Gly 65 70 75 80

Gly Ala Asn Thr Leu Tyr Tyr Ile Lys Asn Leu Asp Gly Gly Asn Lys 85 90 95

Val Ala Asn Val Val Thr Leu Gly Gly Ala Asn Arg Leu Thr Thr Gly 100 105

Lys Ala Pro Pro Gly Thr Asp Pro Asp Gln Lys Ile Leu Tyr Thr Ser 115 120 125

Ile Tyr Ser Ser Asp Asp Met Ile Val Met Asn Tyr Leu Ser Arg Leu 130 135 140

Asp Gly Ala Arg Asn Val Gln Ile His Gly Gly His Ile Gly Leu Leu 145 150 155 160

Tyr Ser Ser Gln Val Tyr Ser Leu Ile Lys Glu Gly Leu Asn Gly Gly 165 170 175

Gly Gln Asn Thr Asn 180

<210> 13

<211> 24

<212> DNA

<213> Artificial Sequence

<220> <223>

Oligonucleotide

<400> 13

cgccag	ggtt ttcccagtca cgac	71914.ST25.txt	24
<210> <211> <212> <213>	14 22 DNA Artificial Sequence		
<220> <223>	Oligonucleotide		
<400> tgacac	14 agga aacagctatg ac		22
<210> <211> <212> <213>	15 28 DNA Artificial Sequence		
<220> <223>	Primer		
<400> ccatga	15 ttac gcatatggct gaacacaa		28
<210> <211> <212> <213>	16 26 DNA Artificial Sequence		
<220> <223>	Primer		
<400> ggagga	16 tcat atgaaatttg taaaaa	į	26
<210> <211> <212> <213>	23		
<220> <223>	Primer		
<400> cccggga	17 atcc attgtccgtt acc		23
<210> <211> <212> <213>	18 25 DNA Artificial Sequence		
<220> <223>	Primer		
<400> cgtcage	18 cgaa ttccgctgaa cacat		25

<210> <211> <212> <213>	19 24 DNA Artificial Sequence	71314.3123.686	
<220> <223>	Primer		
<400> gcggga	19 agga tccgaattcg agct		24
<210> <211> <212> <213>	20 24 DNA Artificial Sequence		
<220> <223>	Primer		
<400> cgccag	20 ggtt ttcccagtca cgac		24
<210> <211> <212> <213>	21 22 DNA Artificial Sequence		
<220> <223>	Primer		
<400> tgacaca	21 agga aacagctatg ac		22
<210> <211> <212> <213>	24		
<220> <223>	Primer		
<400>	22 gcgc ctccgggaac agat		24